

Christopher Jones

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

4,518
citations

304368

22
h-index

476904

29
g-index

30
all docs

30
docs citations

30
times ranked

4638
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Loss in microbial diversity affects nitrogen cycling in soil. ISME Journal, 2013, 7, 1609-1619. | 4.4 | 603 |
| 2 | The unaccounted yet abundant nitrous oxide-reducing microbial community: a potential nitrous oxide sink. ISME Journal, 2013, 7, 417-426. | 4.4 | 529 |
| 3 | Relationship between N-cycling communities and ecosystem functioning in a 50-year-old fertilization experiment. ISME Journal, 2009, 3, 597-605. | 4.4 | 478 |
| 4 | Phylogenetic Analysis of Nitrite, Nitric Oxide, and Nitrous Oxide Respiratory Enzymes Reveal a Complex Evolutionary History for Denitrification. Molecular Biology and Evolution, 2008, 25, 1955-1966. | 3.5 | 424 |
| 5 | Genomics and Ecology of Novel N ₂ O-Reducing Microorganisms. Trends in Microbiology, 2018, 26, 43-55. | 3.5 | 388 |
| 6 | Intergenomic Comparisons Highlight Modularity of the Denitrification Pathway and Underpin the Importance of Community Structure for N ₂ O Emissions. PLoS ONE, 2014, 9, e114118. | 1.1 | 383 |
| 7 | Recently identified microbial guild mediates soil N ₂ O sink capacity. Nature Climate Change, 2014, 4, 801-805. | 8.1 | 364 |
| 8 | Importance of denitrifiers lacking the genes encoding the nitrous oxide reductase for N ₂ O emissions from soil. Global Change Biology, 2011, 17, 1497-1504. | 4.2 | 300 |
| 9 | Ecological and evolutionary factors underlying global and local assembly of denitrifier communities. ISME Journal, 2010, 4, 633-641. | 4.4 | 217 |
| 10 | Soil carbon quality and nitrogen fertilization structure bacterial communities with predictable responses of major bacterial phyla. Applied Soil Ecology, 2014, 84, 62-68. | 2.1 | 162 |
| 11 | Changes in faecal bacteria associated with concentrate and forage-only diets fed to horses in training. Equine Veterinary Journal, 2009, 41, 908-914. | 0.9 | 126 |
| 12 | Phenotypic and genotypic heterogeneity among closely related soil-borne N ₂ - and N ₂ O-producing Bacillus isolates harboring the nosZ gene. FEMS Microbiology Ecology, 2011, 76, 541-552. | 1.3 | 53 |
| 13 | Spatial and phylogeographical analyses of nosZ genes underscore niche differentiation amongst terrestrial N ₂ O reducing communities. Soil Biology and Biochemistry, 2017, 115, 82-91. | 4.2 | 52 |
| 14 | Geospatial variation in co-occurrence networks of nitrifying microbial guilds. Molecular Ecology, 2019, 28, 293-306. | 2.0 | 50 |
| 15 | Soil type overrides plant effect on genetic and enzymatic N ₂ O production potential in arable soils. Soil Biology and Biochemistry, 2016, 100, 125-128. | 4.2 | 47 |
| 16 | Intercropping affects genetic potential for inorganic nitrogen cycling by root-associated microorganisms in Medicago sativa and Dactylis glomerata. Applied Soil Ecology, 2017, 119, 260-266. | 2.1 | 45 |
| 17 | The DNRA-Denitrification Dichotomy Differentiates Nitrogen Transformation Pathways in Mountain Lake Benthic Habitats. Frontiers in Microbiology, 2019, 10, 1229. | 1.5 | 44 |
| 18 | Habitat partitioning of marine benthic denitrifier communities in response to oxygen availability. Environmental Microbiology Reports, 2016, 8, 486-492. | 1.0 | 42 |

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|----|---|-----|-----------|
| 19 | Design and evaluation of primers targeting genes encoding NO-forming nitrite reductases: implications for ecological inference of denitrifying communities. <i>Scientific Reports</i> , 2016, 6, 39208. | 1.6 | 37 |
| 20 | Expression of nirK and nirS genes in two strains of <i>Pseudomonas stutzeri</i> harbouring both types of NO-forming nitrite reductases. <i>Research in Microbiology</i> , 2018, 169, 343-347. | 1.0 | 35 |
| 21 | Soil microbial community analysis using two-dimensional polyacrylamide gel electrophoresis of the bacterial ribosomal internal transcribed spacer regions. <i>Journal of Microbiological Methods</i> , 2007, 69, 256-267. | 0.7 | 25 |
| 22 | Lucerne (<i>Medicago sativa</i>) alters N ₂ O-reducing communities associated with cocksfoot (<i>Dactylis</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 <i>Biology and Biochemistry</i> , 2019, 137, 107547. | 4.2 | 25 |
| 23 | Global Phylogeography of Chitinase Genes in Aquatic Metagenomes. <i>Applied and Environmental Microbiology</i> , 2011, 77, 1101-1106. | 1.4 | 21 |
| 24 | Catch Crop Residues Stimulate N ₂ O Emissions During Spring, Without Affecting the Genetic Potential for Nitrite and N ₂ O Reduction. <i>Frontiers in Microbiology</i> , 2018, 9, 2629. | 1.5 | 17 |
| 25 | Denitrification rates in lake sediments of mountains affected by high atmospheric nitrogen deposition. <i>Scientific Reports</i> , 2020, 10, 3003. | 1.6 | 16 |
| 26 | Assessing costs and benefits of improved soil quality management in remediation projects: A study of an urban site contaminated with PAH and metals. <i>Science of the Total Environment</i> , 2020, 707, 135582. | 3.9 | 13 |
| 27 | Reactive nitrogen restructures and weakens microbial controls of soil N ₂ O emissions. <i>Communications Biology</i> , 2022, 5, 273. | 2.0 | 11 |
| 28 | Minimizing tillage modifies fungal denitrifier communities, increases denitrification rates and enhances the genetic potential for fungal, relative to bacterial, denitrification. <i>Soil Biology and Biochemistry</i> , 2022, 170, 108718. | 4.2 | 6 |
| 29 | Habitat diversity and type govern potential nitrogen loss by denitrification in coastal sediments and differences in ecosystem-level diversities of disparate N ₂ O reducing communities. <i>FEMS Microbiology Ecology</i> , 2020, 96, . | 1.3 | 5 |